

GenCore version: 5.1.4.15-4578  
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GM nucleic acid search, using sw model

Run on: April 6, 2003, 20:06:59 : Search time: 99.5 seconds  
(without alignments)

10182199 Million cells updates/sec

Title: US-08-572-027b-1

Sequence: 1 AAGCTGCAAGTGGACAAAT.....GCTACATCAATTAATCA 1155

Scoring table: IDENTITY: 100  
Gap: 10.0, Gap: 1.0

Search: 59229 seqs, 4292390 residues

Total number of hits satisfying chosen parameters: 118658

Minimum hit seq length: 0

Maximum hit seq length: 200000000

Post-processing: Minimum Match 100%  
Listing first 45 summaries

Database:

1. Published\_Applications\_NA:  
2. 1000 Genomes Project  
3. 1000 Genomes Project  
4. 1000 Genomes Project  
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6. 1000 Genomes Project  
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12. 1000 Genomes Project  
13. 1000 Genomes Project  
14. 1000 Genomes Project

Prod. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	1154	99.9	1155	10	US-09-995-297-1	Sequence 1, Appl1
2	1150.8	99.6	1155	10	US-09-995-297-1	Sequence 2, Appl1
3	1148	99.5	1155	10	US-09-995-297-1	Sequence 3, Appl1
4	1136.4	98.4	1155	10	US-09-995-297-1	Sequence 4, Appl1
5	1136.4	98.4	1155	10	US-09-995-297-1	Sequence 5, Appl1
6	1088.4	94.2	1155	10	US-09-995-297-1	Sequence 6, Appl1
7	1086.8	94.1	1155	10	US-09-995-297-1	Sequence 7, Appl1
8	1085.8	94.1	1155	10	US-09-995-297-1	Sequence 8, Appl1
9	1085.2	94.0	1155	10	US-09-995-297-1	Sequence 9, Appl1
10	720.2	62.4	1155	10	US-09-995-297-1	Sequence 10, Appl1
11	720.2	62.4	1155	10	US-09-995-297-1	Sequence 11, Appl1
12	720.2	62.4	1155	10	US-09-995-297-1	Sequence 12, Appl1
13	566.6	49.1	1111	10	US-09-995-297-1	Sequence 13, Appl1
14	514.4	44.5	1111	10	US-09-995-297-1	Sequence 14, Appl1
15	500.2	43.3	1156	9	US-10-224-446-1	Sequence 15, Appl1
16	473.6	41.0	1156	9	US-10-224-446-1	Sequence 16, Appl1
17	462.4	40.0	1121	9	US-10-224-446-1	Sequence 17, Appl1
18	448.9	38.9	1308	9	US-10-224-446-1	Sequence 18, Appl1
19	448	38.8	1199	9	US-09-981-124-19	Sequence 19, Appl1

Prod. No.	Score	Query Match	Length	DB	ID	Description
20	418.8	34.3	1309	9	US-09-981-124-1	Sequence 20, Appl1
21	409.4	32.4	1308	9	US-09-981-124-1	Sequence 21, Appl1
22	401.8	32.2	765	9	US-10-224-446-2	Sequence 22, Appl1
23	461	41.3	544	10	US-09-885-186-2	Sequence 23, Appl1
24	461	41.3	544	10	US-09-885-186-2	Sequence 24, Appl1
25	429.6	28.5	544	10	US-09-885-186-1	Sequence 25, Appl1
26	429.6	28.5	544	10	US-09-885-186-1	Sequence 26, Appl1
27	429.6	28.5	544	10	US-09-885-186-1	Sequence 27, Appl1
28	408.8	26.7	655	10	US-09-770-149-504	Sequence 28, Appl1
29	275.4	21.6	1301	10	US-09-852-139-1	Sequence 29, Appl1
30	275.4	21.6	1301	10	US-09-852-139-1	Sequence 30, Appl1
31	275.4	21.6	1301	10	US-09-852-139-1	Sequence 31, Appl1
32	194.6	16.8	550	9	US-09-981-124-1	Sequence 32, Appl1
33	164.2	14.4	306	10	US-09-878-574-4920	Sequence 33, Appl1
34	152	13.2	342	10	US-09-878-574-2308	Sequence 34, Appl1
35	145.6	12.6	283	10	US-09-878-574-3638	Sequence 35, Appl1
36	107.4	9.3	177	9	US-09-981-124-1	Sequence 36, Appl1
37	74.4	6.4	245	10	US-09-878-574-3936	Sequence 37, Appl1
38	71.2	6.2	1399	9	US-09-981-124-1	Sequence 38, Appl1
39	64.2	5.6	520	9	US-10-184-444-352	Sequence 39, Appl1
40	57.6	5.0	1301	9	US-10-184-444-352	Sequence 40, Appl1
41	57.6	5.0	1301	9	US-10-184-444-352	Sequence 41, Appl1
42	54.6	4.7	550	9	US-09-878-574-2308	Sequence 42, Appl1
43	54.6	4.7	550	9	US-09-878-574-2308	Sequence 43, Appl1
44	54.6	4.7	550	9	US-09-878-574-2308	Sequence 44, Appl1
45	53.8	4.7	544	9	US-10-184-444-352	Sequence 45, Appl1

## ALIGNMENTS

Result No.	Score	Query Match	Length	DB	ID	Description
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2	1150.8	99.6	1155	10	US-09-995-297-1	Sequence 2, Appl1
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7	1086.8	94.1	1155	10	US-09-995-297-1	Sequence 7, Appl1
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9	1085.2	94.0	1155	10	US-09-995-297-1	Sequence 9, Appl1
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11	720.2	62.4	1155	10	US-09-995-297-1	Sequence 11, Appl1
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14	514.4	44.5	1111	10	US-09-995-297-1	Sequence 14, Appl1
15	500.2	43.3	1156	9	US-10-224-446-1	Sequence 15, Appl1
16	473.6	41.0	1156	9	US-10-224-446-1	Sequence 16, Appl1
17	462.4	40.0	1121	9	US-10-224-446-1	Sequence 17, Appl1
18	448.9	38.9	1308	9	US-10-224-446-1	Sequence 18, Appl1
19	448	38.8	1199	9	US-09-981-124-19	Sequence 19, Appl1



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US-09-995-297-6

Sequence 5, Applicant ID: 15270006207  
; Patent No. US20020092038A1  
; Patent No. US20020092038A1

### STUDYING INTERNATIONAL ADDITIONAL KEYWORD: International

APPLICANT: *Pat, 21000000*

APPL. LEAFHOPPER, COLLEGE

FILE REFERENCE: 07148-072002

CURRENT ACCOUNT NUMBER: 00709095, 277  
CURRENT PAYMENT DATE: 2007 11 27

PRIOR APPLICATION NUMBER: 05/09/128,602

NUMBER OF STUDENTS: 68

: SOFTWARE. Tested for Windows Version 4.0  
: SHQ ID NO 5
$$\frac{1.31 \times 10^4 \text{ g/mol}}{1.155} = 1.13 \times 10^4 \text{ g/mol}$$

CRANISM: brassica napus  
EVALUATE.

1. **FUNCTION:** *to be*  
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LOCATION: (1) ... (1152)  
OTHER INFORMATION: Wild type 1942

US-09-395-297-5

Query Match	94.28;	Score 1088
host local: Similarity	96.49;	Prod No.

Math 1113: Calculus I (M)

QY 1 ATGCTTGCAGTTGAAATGAAATGTTCTGTT

D<sub>b</sub>

1 ATGGCTGCAATCTGGAGAGAACTAATTGTTTCT

61 ATCAAGCGGTAATCTGGACACACGGTCTCG

6)  $\Delta \mu_{\text{max}} = 0.1 \text{ V}$   $\Delta \mu_{\text{max}} = 0.2 \text{ V}$   $\Delta \mu_{\text{max}} = 0.3 \text{ V}$   $\Delta \mu_{\text{max}} = 0.4 \text{ V}$   $\Delta \mu_{\text{max}} = 0.5 \text{ V}$   $\Delta \mu_{\text{max}} = 0.6 \text{ V}$   $\Delta \mu_{\text{max}} = 0.7 \text{ V}$   $\Delta \mu_{\text{max}} = 0.8 \text{ V}$   $\Delta \mu_{\text{max}} = 0.9 \text{ V}$   $\Delta \mu_{\text{max}} = 1.0 \text{ V}$

DOI: 10.1002/anie.200500016

**QY**    |Z| CACGGAGCGTCAATGCTCCGATCCTCCG  
|| || || || || || || || || || || || || || || ||

**D**b    : 21   CCGACCTGAAATGGTTCAGGTTCCGCTCCTG

181 AITAMOTUWONGTAWANINUTAN

[81]  $Al_2(AiAlO_2)_2$  [57] [58] [10] [11] [7] [16] [17] [18] [19] [20] [21] [22] [23] [24] [25] [26] [27] [28] [29] [30] [31] [32] [33] [34] [35] [36] [37] [38] [39] [40] [41] [42] [43] [44] [45] [46] [47] [48] [49] [50] [51] [52] [53] [54] [55] [56] [59] [60] [61] [62] [63] [64] [65] [66] [67] [68] [69] [70] [71] [72] [73] [74] [75] [76] [77] [78] [79] [80] [81] [82] [83] [84] [85] [86] [87] [88] [89] [90] [91] [92] [93] [94] [95] [96] [97] [98] [99] [100] [101] [102] [103] [104] [105] [106] [107] [108] [109] [110] [111] [112] [113] [114] [115] [116] [117] [118] [119] [120] [121] [122] [123] [124] [125] [126] [127] [128] [129] [130] [131] [132] [133] [134] [135] [136] [137] [138] [139] [140] [141] [142] [143] [144] [145] [146] [147] [148] [149] [150] [151] [152] [153] [154] [155] [156] [157] [158] [159] [160] [161] [162] [163] [164] [165] [166] [167] [168] [169] [170] [171] [172] [173] [174] [175] [176] [177] [178] [179] [180] [181] [182] [183] [184] [185] [186] [187] [188] [189] [190] [191] [192] [193] [194] [195] [196] [197] [198] [199] [200] [201] [202] [203] [204] [205] [206] [207] [208] [209] [210] [211] [212] [213] [214] [215] [216] [217] [218] [219] [220] [221] [222] [223] [224] [225] [226] [227] [228] [229] [230] [231] [232] [233] [234] [235] [236] [237] [238] [239] [240] [241] [242] [243] [244] [245] [246] [247] [248] [249] [250] [251] [252] [253] [254] [255] [256] [257] [258] [259] [260] [261] [262] [263] [264] [265] [266] [267] [268] [269] [270] [271] [272] [273] [274] [275] [276] [277] [278] [279] [280] [281] [282] [283] [284] [285] [286] [287] [288] [289] [290] [291] [292] [293] [294] [295] [296] [297] [298] [299] [300] [301] [302] [303] [304] [305] [306] [307] [308] [309] [310] [311] [312] [313] [314] [315] [316] [317] [318] [319] [320] [321] [322] [323] [324] [325] [326] [327] [328] [329] [330] [331] [332] [333] [334] [335] [336] [337] [338] [339] [340] [341] [342] [343] [344] [345] [346] [347] [348] [349] [350] [351] [352] [353] [354] [355] [356] [357] [358] [359] [360] [361] [362] [363] [364] [365] [366] [367] [368] [369] [370] [371] [372] [373] [374] [375] [376] [377] [378] [379] [380] [381] [382] [383] [384] [385] [386] [387] [388] [389] [390] [391] [392] [393] [394] [395] [396] [397] [398] [399] [400] [401] [402] [403] [404] [405] [406] [407] [408] [409] [410] [411] [412] [413] [414] [415] [416] [417] [418] [419] [420] [421] [422] [423] [424] [425] [426] [427] [428] [429] [430] [431] [432] [433] [434] [435] [436] [437] [438] [439] [440] [441] [442] [443] [444] [445] [446] [447] [448] [449] [450] [451] [452] [453] [454] [455] [456] [457] [458] [459] [460] [461] [462] [463] [464] [465] [466] [467] [468] [469] [470] [471] [472] [473] [474] [475] [476] [477] [478] [479] [480] [481] [482] [483] [484] [485] [486] [487] [488] [489] [490] [491] [492] [493] [494] [495] [496] [497] [498] [499] [500] [501] [502] [503] [504] [505] [506] [507] [508] [509] [510] [511] [512] [513] [514] [515] [516] [517] [518] [519] [520] [521] [522] [523] [524] [525] [526] [527] [528] [529] [530] [531] [532] [533] [534] [535] [536] [537] [538] [539] [540] [541] [542] [543] [544] [545] [546] [547] [548] [549] [550] [551] [552] [553] [554] [555] [556] [557] [558] [559] [560] [561] [562] [563] [564] [565] [566] [567] [568] [569] [570] [571] [572] [573] [574] [575] [576] [577] [578] [579] [580] [581] [582] [583] [584] [585] [586] [587] [588] [589] [590] [591] [592] [593] [594] [595] [596] [597] [598] [599] [600] [601] [602] [603] [604] [605] [606] [607] [608] [609] [610] [611] [612] [613] [614] [615] [616] [617] [618] [619] [620] [621] [622] [623] [624] [625] [626] [627] [628] [629] [630] [631] [632] [633] [634] [635] [636] [637] [638] [639] [640] [641] [642] [643] [644] [645] [646] [647] [648] [649] [650] [651] [652] [653] [654] [655] [656] [657] [658] [659] [660] [661] [662] [663] [664] [665] [666] [667] [668] [669] [670] [671] [672] [673] [674] [675] [676] [677] [678] [679] [680] [681] [682] [683] [684] [685] [686] [687] [688] [689] [690] [691] [692] [693] [694] [695] [696] [697] [698] [699] [700] [701] [702] [703] [704] [705] [706] [707] [708] [709] [710] [711] [712] [713] [714] [715] [716] [717] [718] [719] [720] [721] [722] [723] [724] [725] [726] [727] [728] [729] [730] [731] [732] [733] [734] [735] [736] [737] [738] [739] [740] [741] [742] [743] [744] [745] [746] [747] [748] [749] [750] [751] [752] [753] [754] [755] [756] [757] [758] [759] [760] [761] [762] [763] [764] [765] [766] [767] [768] [769] [770] [771] [772] [773] [774] [775] [776] [777] [778] [779] [780] [781] [782] [783] [784] [785] [786] [787] [788] [789] [790] [791] [792] [793] [794] [795] [796] [797] [798] [799] [800] [801] [802] [803] [804] [805] [806] [807] [808] [809] [810] [811] [812] [813] [814] [815] [816] [817] [818] [819] [820] [821] [822] [823] [824] [825] [826] [827] [828] [829] [830] [831] [832] [833] [834] [835] [836] [837] [838] [839] [840] [841] [842] [843] [844] [8

**QY** 441 670 10<sup>8</sup> 1A' (100%) 100% 1A' (100%) 100%

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by [10] "RESEARCH AND ANALYSIS OF THE EFFECTS OF THE 1997-1998 EL NIÑO ON THE ECONOMY OF THE UNITED STATES."

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**DE** SOCIETATE ALMA MATER DE GRADUATOS DE LA UNIV.

[illegible]

**Pd**    **69**    **ACCGTGGCTGATGTTCAATTGTTTCCTTC**

421 CHINESE JOURNAL OF ANATOMY

**Db** 421 CACCACTTAAAGTTGGG







[illegible]

CY	11 48	TATACATAAATAAATAA	1155
DB	15 69	TATACATAAATAAATAA	1556
 RESULTS 11 US-09 885-189 3 Sequence by Application 55/00885189 Patent No. US2002015169A1  GENERAL INFORMATION: APPLICANT: CHRIS SOMERVILLE APPLICANT: PETER JORSEN APPLICATION: FRANK VAN DE LIND TITLE OF INVENTION: Production of Hyfluxated Parly Acids In TITLE OF INVENTION: Synthetically Modified Plurals NUMBER OF CLAIMS: 15  CORRESPONDENCE ADDRESS: ADDRESSEE: PHILIPPO MADISON & SUTRO, LLP STREET: 1100 NEW YORK AVENUE, N.W. CITY: WASHINGTON STATE: D.C. COUNTRY: USA ZIP: 20005-5018  COMPUTER READABLE FORM: MEDIUM TYPE: Diskette, 4.50 Inch OPERATING SYSTEM: MS-DOS/MS-DOS SOFTWARE: Word Perfect 5.1  CURRENT APPLICATION DATA: ALTERNATE # ENTERED: 7/7/97 139 FILING DATE: 21 June 2001 CLASSIFICATION:  PRIOR APPLICATION DATA: APPLICATION NUMBER: US/582/597-4140 FILING DATE: February 6, 1996 PARENT APPLICATION IN DATA: APPLICATION NUMBER: 06/730,962 FILING DATE: September 20, 1995  PRIOR APPLICATION DATA: APPLICATION NUMBER: 06/520,984 FILING DATE: October 11, 1994  ENTER ALTERNATIVE DATA: APPLICATION NUMBER: 08/714,596 FILING DATE: September 4, 1994  INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS: LENGTH: 1855 nucleotides TYPE: nucleotide STRATEGY: single TOPOLOGY: linear US-09-885-189 4			
Query Match	62.4%	Score 720.27	Dn 100 Length 1855
Best Local Similarity	77.5%	Prod. No. 1.20E+227	
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DB	402	ATGGGTTTTTGTTGAAAGAAATTAAGATCTGTCTTTGAAGAAGATTAAGATTAAGACT	463
DB	492	CTAAAGGAGGAGGATCGAAAGAGGAGATTAAGATCTGTCTTTGAAGAAGATTAAGACT	522
CY	121	CGAACGTATCTTTGAAGGCTTCAGGATCTGTCTTTGTTGTTGTTGTTGTTGTTGTTGTT	189
DB	522	CGCAAGAGTTCTTGAAGAGCTTATAGAGGATCTGTCTTTGTTGTTGTTGTTGTTGTTGTT	588
CY	181	ATCAACAATGCTTCTTCTTCAATCAATGCTTCTTCTTCAATCAATGCTTCTTCTTCAATCA	244
DB	592	ACTTATATCTCTTCTTCTTCAATCAATGCTTCTTCTTCAATCAATGCTTCTTCTTCAATCA	644
CY	241	CTCATCTCAATCTGCTGAGGCTCTTCACTGAGGCTCTTCACTGAGGCTCTTCACTGAGGCT	297



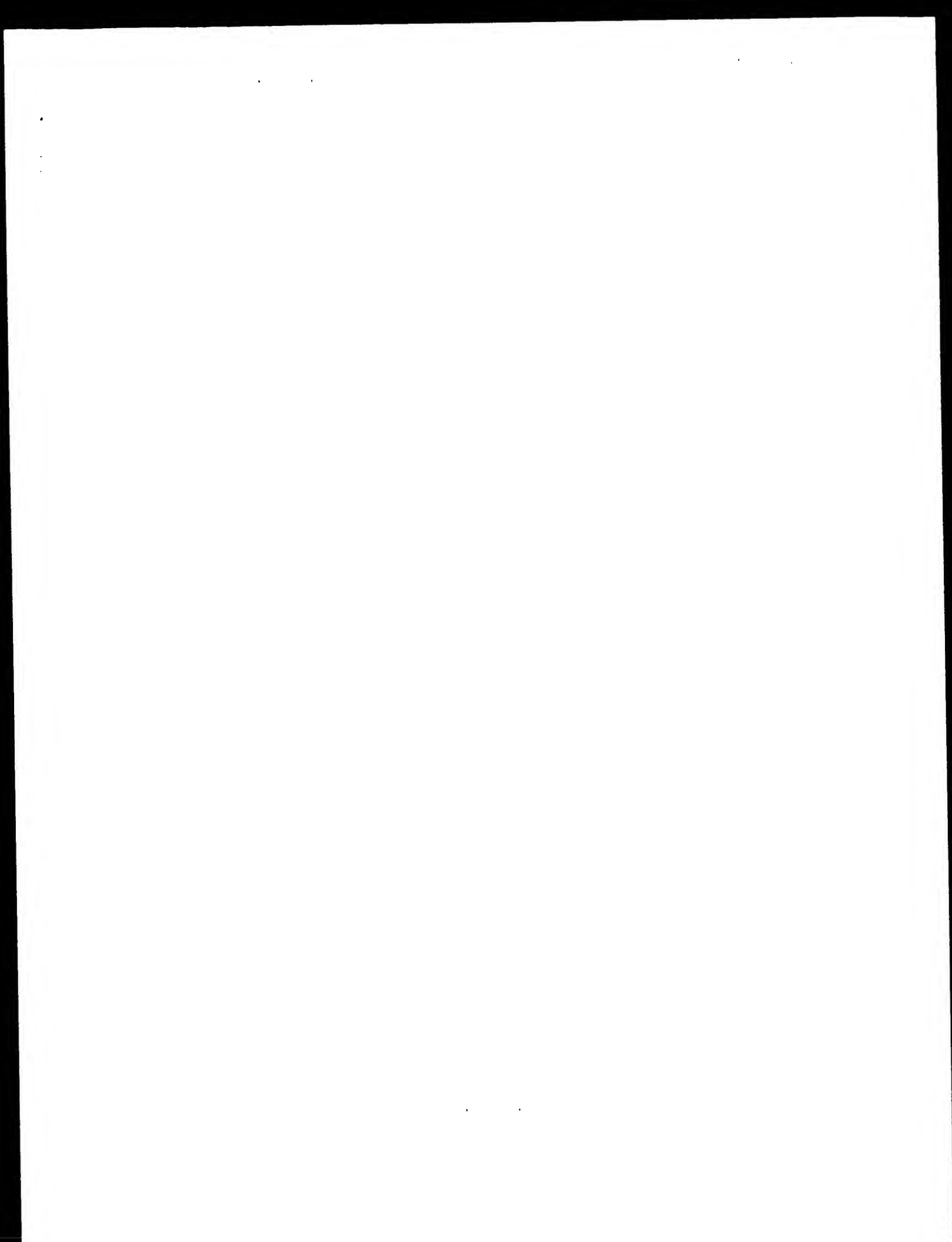






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UY 290 TAAACGAGGCTCTGAGTATATCCAGAAATGAGGAGCAAGGCTTCAGACATATAGT 349  
DB 304 AACTTGTATATGCTTTATGAGCAAGAGGAGGAGCAAGGCTTCAGACATATAGT 367  
UY 353 GAGTTACAGACATGAGGCTGATCTTACCTGCTTCTGCTGCTTATCTTCT 406  
DB 368 GAGTAAAGACAGATGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 427  
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DB 428 GGAAGTACAGTATGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 487  
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DB 488 ATCCGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 547  
-UY 530 GAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 589  
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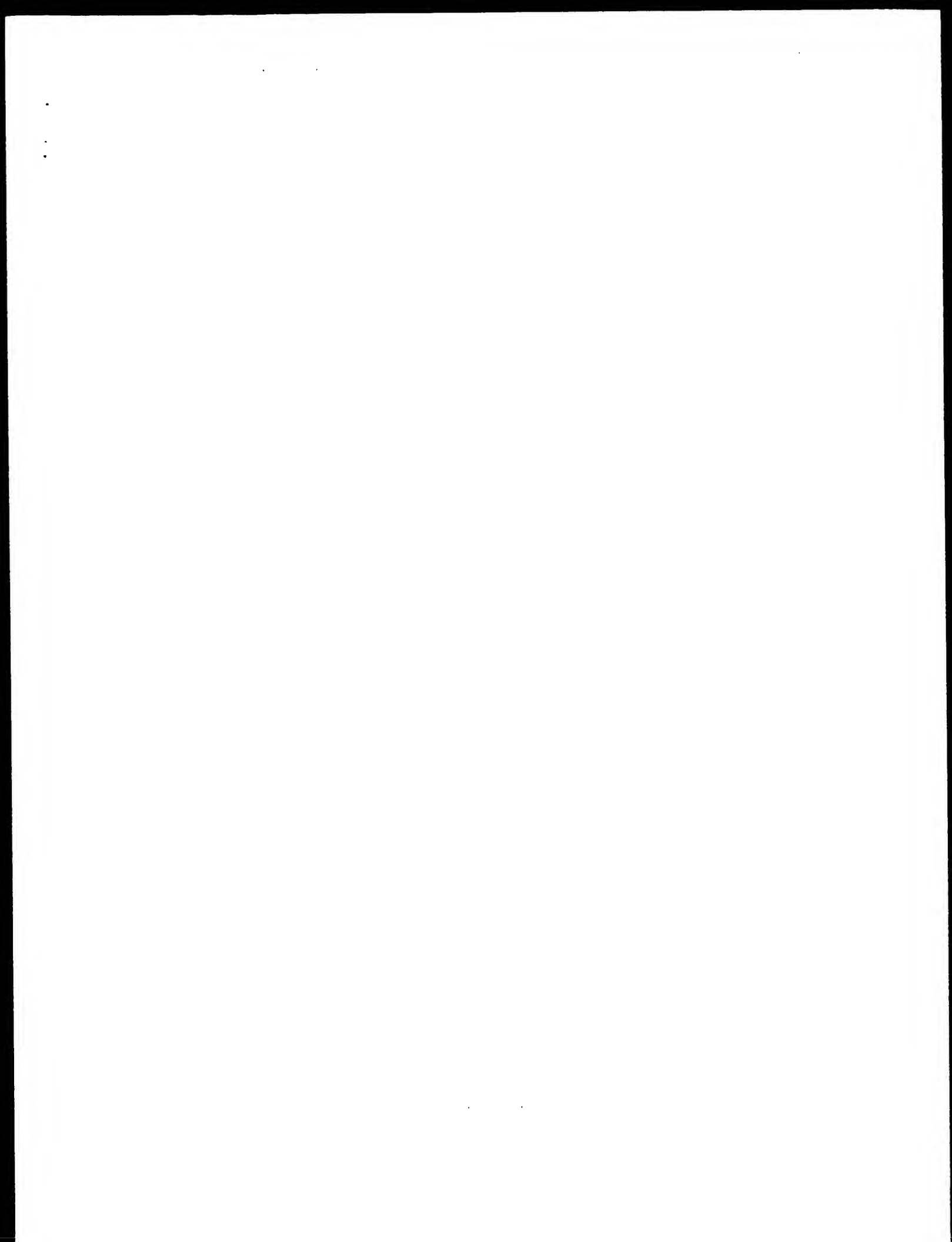


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RESULT 10
US-09-128-602B-9
: Sequence 9, Application US/09128602B
: Patent No. 6414223
: GENERAL INFORMATION:
: APPLICANT: Kodali, Dharna
: APPLICANT: Fan, Zhongqiong
: APPLICANT: DeBonte, Lorin P.
: TITLE OF INVENTION: PLANTS, SEEDS AND OILS HAVING AN ELEVATED TOTAL MONOUNSATURATED
: TITLE OF INVENTION: FATTY ACID CONTENT
: FILE REFERENCE: 07148-072001
: PRESENT APPLICATION NUMBER: US/09-128-602B
: CURRENT FILING DATE: 1998-08-03
: NUMBER OF SEQ ID NOS: 68
: SOFTWARE: FASTSEQ for Windows Version 4.0
: SEQ ID NO: 9
: LENGTH: 1155
: TYPE: DNA
: ORGANISM: Brassica napus
: FEATURE:
: NAME/KEY: CDS
: LOCATION: (1)...(1152)
: US-09-128-602B-9

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## RESULT 15

US-08-572-027b-5.rml

Sequence 3: Application US/086756F08

Patient No. 5850826

GENERAL INFORMATION:

APPLICANT: Labco, Inc. et al.

TITLE OF INVENTION: CANOLA OIL HAVING INCREASED OLEIC ACID AND

TITLE OF INVENTION: DECREASED LINOLENIC ACID CONTENT

NUMBER OF SEQUENCES: 6

COPIES/SEQUENCE ADDED:

ADDRESSER: Fish &amp; Richardson, P.C., P.A.

STREET: 60 South Sixth Street, Suite 3300

CITY: Minneapolis

STATE: MN

COUNTRY: USA

ZIP: 55402

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

OPERATING SYSTEM: PC/MS-DOS

SOFTWARE: Patient Release #1.0, Version #1.00

APPLICATION NUMBER: US/08/675,650B

FILING DATE: 03-10-1996

CLASSIFICATION: 800

ATTORNEY/AGENT INFORMATION:

NAME: Lundquist, Ronald G.

REGISTERED IN NUMBER: 57,875

REFERENCE TO PCT NUMBER: 97/49742001

TELEPHONE: 612/335-5070

TELEFAX: 612/335-5070

INFORMATION FOR SEQ ID NO: 3:

SEQUENCE CHARACTERISTICS:

LENGTH: 1155 base pairs  
 TYPE: nucleic acid  
 STRANDNESS: single  
 TOPOLOGY: linear  
 MOLECULE TYPE: RNA  
 HYPOTHETICAL: NO  
 ARTI SEQUENCE: NO  
 ORIGINAL SOURCE:  
 ORGANISM: Brassica napus  
 IMMEDIATE SOURCE:  
 CLONING: JMS129  
 FEATURES:  
 OTHER INFORMATION: mutation of nucleotide 115 of the 1155  
 US-08-572-027b-5.rml

Query Match: 99.08% Score: 114.44 DB: 27 Length: 1155  
 Host: Local Similarity: 99.08% Prod. No. 1.5e-294  
 Ratios: 1144 Conserved: 50 Mismatches: 60 Indels: 00 Gaps: 00

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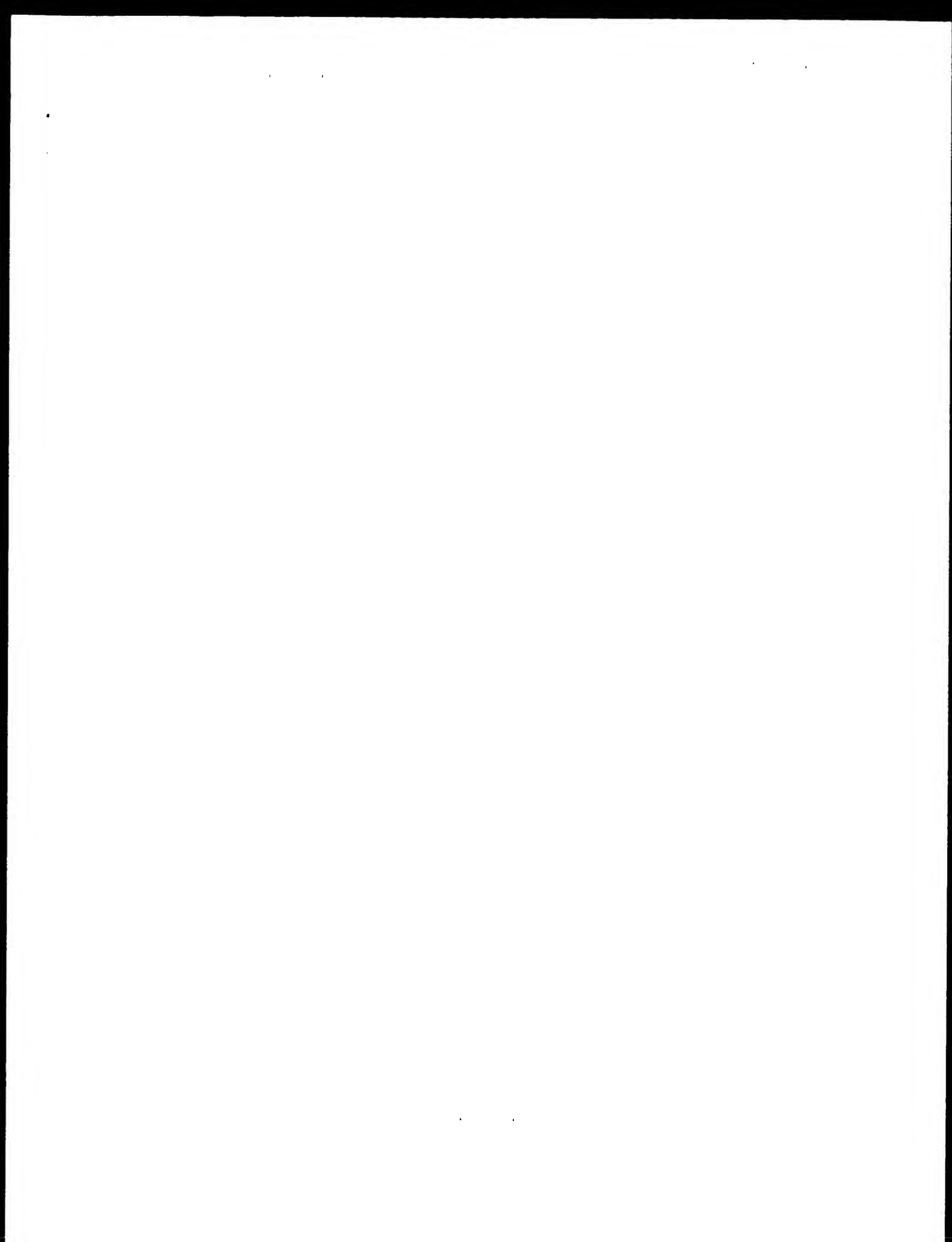








































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 DB 841 GTGATTCGATCGTTCTCTGCTGCTGATCACTTACTTCGACACACGACATCTTCCTG 841  
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## RESULT 13

US-08-728-027a-5

US-08-728-027a-5

GENERAL INFORMATION:

APPLICANT: KIM, GEORGE

TITLE OF INVENTION: GENES FOR METABOLIC

TITLE OF INVENTION: FATTY ACID ELISA 12

TITLE OF INVENTION: DESATURASES AND RELATED

TITLE OF INVENTION: ENZYMES FROM PLANTS

NUMBER OF SEQUENCES: 16

CORRESPONDING AGENCIES:

ADDRESSEE: E. L. POINT DE NEUMERS

ADDRESS: 1007 MARKET STREET

CITY: WILMINGTON

STATE: DELAWARE

COUNTRY: U.S.A.

ZIP: 19898

COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: MICROSOFT WINDOWS 4.1

SOFTWARE: MICROSOFT WORD 6.0

CURRENT APPLICATION DATA:

FILING DATE: 03/29/99

CLASSIFICATION: 435

PRIORITY APPLICATION DATA:

FILING DATE: NOVEMBER 17, 1992

ATTORNEY: ARTHUR J. HANLEY

NAME: SINGELL, BARBARA G.

REGISTRATION NUMBER: 50,764

REFERENCE/DOCKET NUMBER: BR 1043-C

TELECOMMUNICATION INFORMATION:

TELEPHONE: (302) 992-4941

TELEFAX: (302) 774-0164

TELEX: 835420

INFORMATION FOR SEQ ID NO. 5:

SEQUENCE CHARACTERISTICS:

LENGTH: 1156 base pairs

TYPE: nucleic acid

STRANDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: DNA  
 HYPOTHETICAL: NO  
 ANTI-SENSE: NO  
 ORIGINAL SOURCE:  
 ORGANISM: Brassica napus  
 FEATURE:  
 OTHER INFORMATION: wild type F form.  
 US-08-728-027a-5

Query Match 99.88% Score 1162.4 DB 11 Length 1155  
 Best Local Similarity 99.88% Pct. No. 1162-266  
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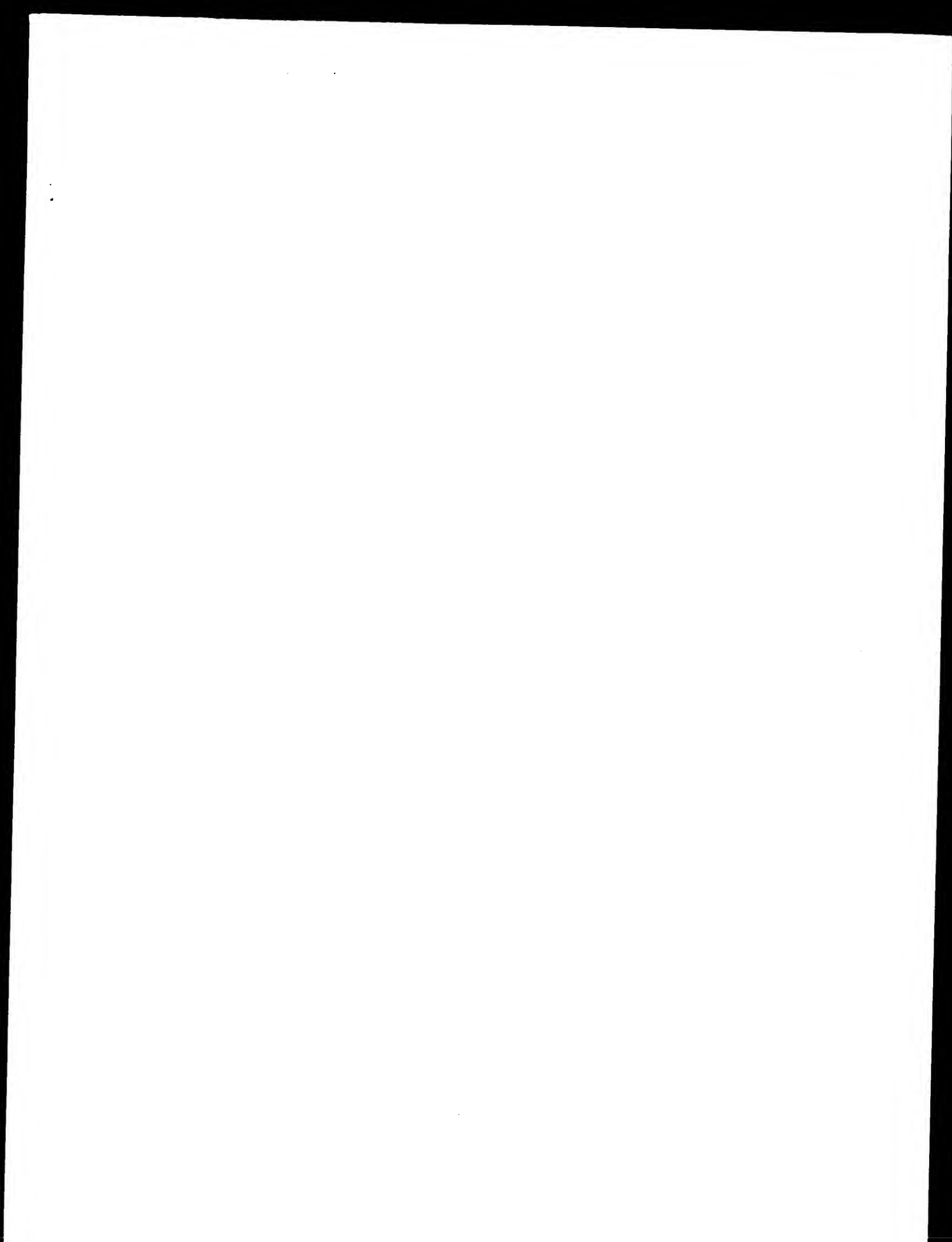






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